

digested M13mp19. Thus, as in Example I, one set of oligonucleotides begins with the addition of A followed by nine rounds of split and mix synthesis wherein the oligonucleotide is extended subunit-wise by 3'-phosphoramidite derivatized 4-mers corresponding to the subunits of Table I. The synthesis is then completed with the nucleotide-by-nucleotide addition of one half of the Sma I recognition site (GGG), two C's, and a 5'-monophosphate, e.g. via the Phosphate-ON reagent available from Clontech Laboratories (Palo Alto, CA). The other set of oligonucleotides begins with the addition of three C's (portion of the Sma I recognition site) and two G's, followed by nine rounds of split and mix synthesis wherein the oligonucleotide is extended by 3'-phosphoramidite derivatized 4-mers corresponding to the complements of the subunits of Table I. Synthesis is completed by the nucleotide-by-nucleotide addition of the Hind III recognition site and a 5'-monophosphate. After separation from the synthesis supports the oligonucleotides are mixed under conditions that permit formation of the following duplexes ([SEQ ID NO:18](#)):

5' -pGGGCC(w<sub>1</sub>)(w<sub>1</sub>)(w<sub>1</sub>)(w<sub>1</sub>)(w<sub>1</sub>)(w<sub>1</sub>)(w<sub>1</sub>)(w<sub>1</sub>)(w<sub>1</sub>)(w<sub>1</sub>)A  
 CCCGG(\*\*)(\*\*)(\*\*)(\*\*)(\*\*)(\*\*)(\*\*)(\*\*)(\*\*)(\*\*) TTCGAP-5'

The mixture of duplexes is then ligated into a Sma I/Hind III-digested M13mp19. A repertoire of tag complements are synthesized on CPG microparticles as described above.<sup>26</sup>

5. Please amend the paragraph in column 25, lines 61-67, as follows:

"After hybridization and ligation, as described in Example 1, the loaded microparticles are treated with Fok I to produce a 4-nucleotide protruding strand of a predetermined sequence. A 10:1 mixture (probe 1:probe 2) of the following probes (SEQ ID NO:3, SEQ ID NO:8[~~, SEQ ID NO:9,~~ and SEQ ID NO:10]) are ligated to the polynucleotides on microparticles."

**IN THE SEQUENCE LISTING:**

From columns 29 and 30, line 30, to columns 35 and 36, line 14, please delete the Sequence Listing and replace it with the following:

- Sequence Listing

<110> Brenner, Sydney  
<120> Compositons for Sorting Polynucleotides  
<130> 802-04RE  
<140> US 09/366,081  
<141> 1999-08-02  
<150> US 08/484,712  
<151> 1995-06-07  
<150> US 08/358,810  
<151> 1994-12-19  
<150> US 08/322,348  
<151> 1994-10-13  
<160> 19  
<170> Microsoft Word97

*Sub*  
*D*

<210> 1  
<211> 38  
<212> DNA  
<213> Artificial Sequence  
<220>  
<223> Segment of vector.  
<400> 1  
gaggatgcct ttatggatcc actcgagatc ccaatcca 38

<210> 2  
<211> 26  
<212> DNA  
<213> Artificial Sequence  
<220>  
<223> Adaptor.  
<400> 2  
aattcggatg atgcatgcat cgaccc 26

<210> 3  
<211> 14  
<212> DNA  
<213> Artificial Sequence  
<220>  
<223> Adaptor.  
<400> 3  
tcgagtcatc cgat 14

*C*  
*le*

<210> 4  
<211> 39  
<212> DNA  
<213> Artificial Sequence  
<220>  
<223> Tag complement linked to solid phase support.  
<400> 4  
ddddddddd dddddd dddddd dddddd tgg 39

<210> 5  
<211> 68  
<212> DNA  
<213> Artificial Sequence  
<220>  
<223> Primer for synthesis of first strand of cDNA. Primer contains tag sequence.  
<400> 5  
ctagtcgacc ahhhhhhhhh hhhhhhhhhh hhhhhhhhhh hhhhhhhhggt 50  
tttttttttt tttttttt 68

<210> 6  
<211> 11  
<212> DNA  
<213> Artificial Sequence  
<220>  
<221> any of a, c, g, t, or u at indicated position  
<222> 1, 9-11

*Sub 2*

<223> a, c, g, t, or u  
<400> 6

nrrgatcynn n

11

<210> 7  
<211> 22  
<212> DNA  
<213> Artificial Sequence  
<220>  
<223> Adaptor.  
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gggtcgatgc atgcatcatc cg

22

<210> 8  
<211> 10  
<212> DNA  
<213> Artificial Sequence  
<220>  
<223> Adaptor.  
<400> 8

atcggatgac

10

<210> 9  
<211> 43  
<212> DNA  
<213> Artificial Sequence  
<220>  
<223> Adaptor containing oligonucleotide tag.  
<400> 9

tcgacchhhh hhhhhhhhhh hhhhhhhhhh hhhhhhhhhh hha

43

<210> 10  
<211> 43  
<212> DNA  
<213> Artificial Sequence  
<220>  
<223> Adaptor containing oligonucleotide tag.  
<400> 10

tcgacchhhh hhhhhhhhhh hhhhhhhhhh hhhhhhhhhh hha

43

<210> 11  
<211> 16  
<212> DNA  
<213> Artificial Sequence  
<220>  
<223> Adaptor.  
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atcggatgac atcaac

16

<210> 12  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
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<221> any of a, c, g, t, or u at indicated position

*Sub 2*

<222> 1-3  
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<400> 12  
nnnagttgat gtcatccgat

<210> 13  
<211> 20  
<212> DNA  
<213> Artificial Sequence  
<220>

<221> any of a, c, g, t, or u at indicated position  
<222> 1-3  
<223> a, c, g, t, or u  
<400> 13  
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20

*Sub 2*

<210> 14  
<211> 20  
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<213> Artificial Sequence  
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<221> any of a, c, g, t, or u at indicated position  
<222> 1-3  
<223> a, c, g, t, or u  
<400> 14  
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20

<210> 15  
<211> 20  
<212> DNA  
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<223> a, c, g, t, or u  
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20

<210> 16  
<211> 37  
<212> DNA  
<213> Artificial Sequence  
<220>

<221> any of a, c, g, t, or u at indicated position  
<222> 1-5, 10-23, 25-37  
<223> a, c, g, t, or u  
<400> 16  
nnnnnggatg nnnnnnnnnn nnntnnnnnn nnnnnnnn

37

<210> 17  
<211> 43  
<212> DNA  
<213> Artificial Sequence  
<220>

<223> Adaptor containing oligonucleotide tag.